



## RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: 09/993,808A  
Source: DIPF  
Date Processed by STIC: 3/13/03

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2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY  
FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.  
PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)  
PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

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1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
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OIPE

Does Not Comply  
Corrected Diskette Needed

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/993,808A

DATE: 03/13/2003  
TIME: 14:53:48

Input Set : A:\1146.SEQLIST.TXT  
Output Set: N:\CRF4\03132003\I993808A.raw

4 <110> APPLICANT: Gordon-Kamm, William  
5       Lowe, Keith  
6       Sun, Yuejin  
7       Dilkes, Brian  
8       Larkins, Brian  
11 <120> TITLE OF INVENTION: Cell Cycle Nucleic Acids, Polypeptides,  
12       and Uses Thereof  
14 <130> FILE REFERENCE: 1146  
C--> 16 <140> CURRENT APPLICATION NUMBER: US/09/993,808A  
C--> 16 <141> CURRENT FILING DATE: 2003-03-13  
16 <160> NUMBER OF SEQ ID NOS: 6  
18 <170> SOFTWARE: FastSEQ for Windows Version 3.0

## ERRORED SEQUENCES

109 <210> SEQ ID NO: 2  
110 <211> LENGTH: 102 — found 256  
111 <212> TYPE: PRT  
112 <213> ORGANISM: zea mays  
114 <220> FEATURE:  
115 <221> NAME/KEY: VARIANT  
116 <222> LOCATION: (1)...(256)  
117 <223> OTHER INFORMATION: Xaa = Any Amino Acid  
119 <400> SEQUENCE: 2  
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121       1                       5                       10                       15  
122 Ala Ala Val Glu Val Thr Gln Val Val Gly Val Arg Thr Arg Ser Arg  
123       20                       25                       30  
124 Ser Ala Ala Ala Thr Gly Gly Val Ala Lys Val Ala Pro Arg Arg Lys  
125       35                       40                       45  
126 Arg Ala Pro Ala Gly Glu Pro Ala Ala Ala Val Ser Ala Gly Gly Asp  
127       50                       55                       60  
128 Gly Gly Ser Cys Tyr Ile His Leu Arg Ser Arg Met Leu Phe Met Ala  
129       65                       70                       75                       80  
130 Pro Pro Gln Pro Gln Pro Ser Val Asp Ser Val Pro Thr Pro Val Glu  
131       85                       90                       95  
132 Ala Ala Asp Gly Ala Ala Gly Gln Gln Gly Ala Ala Leu Ala Ala Gly  
133       100                       105                       110  
134 Leu Ser Arg Cys Ser Ser Thr Ala Ser Ser Val Asn Leu Gly Leu Gly  
135       115                       120                       125  
136 Gly Gln Arg Gly Ser His Thr Cys Arg Ser Tyr Asp Ala Ala Glu Ala  
137       130                       135                       140

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138 Gly Gly Asp His Val Leu Val Asp Val Ser Ala Ala Ser Asn Ser Gly
139 145 150 155 160
140 Ser Gly Pro Asp Arg Glu Arg Arg Glu Thr Thr Pro Ser Ser Arg Ala
141 165 170 175
142 His Gly Glu Leu Ser Asp Leu Glu Ser Asp Leu Ala Gly His Lys Thr
143 180 185 190
144 Gly Pro Ser Leu Pro Ala Ala Thr Pro Ala Ala Glu Leu Ile Val Pro
145 195 200 205
146 Pro Ala His Glu Ile Gln Glu Phe Phe Ala Ala Ala Glu Ala Ala Gln
147 210 215 220
148 Ala Lys Arg Phe Ala Ser Lys Tyr Asn Phe Asp Phe Val Arg Gly Val
149 225 230 235 240
150 Pro Leu Asp Ala Gly Gly Arg Phe Glu Trp Ala Pro Val Val Ser Ile
E--> 151 245 250 255

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**VERIFICATION SUMMARY**

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Output Set: N:\CRF4\03132003\I993808A.raw

L:16 M:270 C: Current Application Number differs, Replaced Current Application No  
L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:151 M:252 E: No. of Seq. differs, <211> LENGTH:Input:702 Found:256 SEQ:2  
L:299 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:5  
L:299 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:512  
L:354 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:96